

**NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING
NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES**

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CAR §1.821 - §1.825 for the following reason(s):

- 1. This application clearly fails to comply with the requirements of 37 CAR §1.821 - §1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990, and at 55 FR 18230, May 1, 1990.
- 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 CAR §1.821(c).
- 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 CAR §1.821(e).
- 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 CAR §1.822 and/or §1.823, as indicated on the attached copy of the marked-up "Raw Sequence Listing".
- 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A substitute computer readable form must be submitted as required by 37 CAR §1.825(d).
- 6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CAR §1.821(e).
- 7. Other: _____

APPLICANT MUST PROVIDE:

- An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
- An initial or substitute paper copy of the "Sequence Listing", as were as an amendment directing its entry into the specification.
- A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 CAR §1.821(e) or §1.821(f) or §1.821(g) or §1.825(b) or §1.825(d).

FOR QUESTIONS REGARDING COMPLIANCE WITH THESE REQUIREMENTS, PLEASE CONTACT:

- For Rules Interpretation, call (703) 308-1123
- For CRF Submission help, call (703)308-4212
- For Patentin Software help, call (703) 557-0400

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE.



RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/446,677

DATE: 09/06/2000
TIME: 11:01:37

Input Set : A:\Birkell.txt
Output Set: N:\CRF3\09062000\I446677.raw

Does Not Comply
Corrected Diskette Needed

2, 4, 6
See pp.

SEQUENCE LISTING

3 (1) GENERAL INFORMATION:
5 (i) APPLICANT: BIRKELUND, Svend
6 CHRISTIANSEN, Gunna
7 HEBSGARD PEDERSEN, Anna-Sofie
8 MYGIND, Per
9 KNUDSEN, Katrine
11 (ii) TITLE OF INVENTION: SURFACE EXPOSED PROTEINS FROM CHLAMYDIA
12 PNEUMONIAE
14 (iii) NUMBER OF SEQUENCES: 30
16 (iv) CORRESPONDENCE ADDRESS:
17 (A) ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
18 (B) STREET: 624 Ninth Street, N.W., Suite 300
19 (C) CITY: Washington
20 (D) STATE: D.C.
21 (E) COUNTRY: USA
22 (F) ZIP: 20001
24 (v) COMPUTER READABLE FORM:
25 (A) MEDIUM TYPE: Floppy disk
26 (B) COMPUTER: IBM PC compatible
27 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
28 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
30 (vi) CURRENT APPLICATION DATA:
31 (A) APPLICATION NUMBER: US/09/446,677
32 (B) FILING DATE: 24-Mar-2000
C--> 33 (vii) PRIOR APPLICATION DATA:
34 (A) APPLICATION NUMBER: PCT/DK98/00266
35 (B) FILING DATE: 19-JUN-1998
36 (A) APPLICATION NUMBER: DK 0744/97
37 (B) FILING DATE: 23-JUN-1997
40 (viii) ATTORNEY/AGENT INFORMATION:
41 (A) NAME: COOPER, Iver P.
42 (B) REGISTRATION NUMBER: 28,005
43 (C) REFERENCE/DOCKET NUMBER: BIRKELUND=1
44 (ix) TELECOMMUNICATION INFORMATION:
45 (A) TELEPHONE: 202-628-5197
46 (B) TELEFAX: 202-737-3528
47

ERRORED SEQUENCES

498 (2) INFORMATION FOR SEQ ID NO: 3:
500 (i) SEQUENCE CHARACTERISTICS:
501 (A) LENGTH: 2815 base pairs
502 (B) TYPE: nucleic acid
503 (C) STRANDEDNESS: single
504 (D) TOPOLOGY: linear

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/446,677

DATE: 09/06/2000
TIME: 11:01:37

Input Set : A:\Birkell.txt
Output Set: N:\CRF3\09062000\I446677.raw

Enter "hard return"
here to correct.

506 (ii) MOLECULE TYPE: Genomic DNA
508 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

E--> 510 ATGAAATCGC AATTTCCCTG GTTAGTGCTC TCTTCGACAT TGCCATGTTT TACTAGTTGT 60 TCCACTGTTT TTGCTGCAAC TGCTGAAAAT ATAC
 511 ACTAACACAG GCACCTATAAC TCCTAAAAAT ACAGACTACTG GAATGAGACTA TACTCTGACA 180
 512 GGAGATATAA CTCTGCAAAA CCTTGGGAT TCGGCAGCTT TAACGAAGGG TTGTTTTCT 240
 513 GACACTACGG AATCTTAAAG CTTTCCGGT AAGGGGACT CACTTCTTT TTTAAATATT 300
 514 AAGTCTAGT CTGAGGGCGC AGACATTCT GTTACAACAG 360
 515 GGATTTCTGA GTCTTACTTT CTTAGCGGCC CCATCATCGG TAATCACAAC CCCCTCAAGGA 420
 516 AAAGGTCAG TAAATGTTGG AGGGATCTT ACATTTGATA ACAATGGAAC TATTTTATT 480
 517 AAACAAGATT ACTCTGAGGA AAATGGCGGA GCCATTCTCA CCAAGAATCT TTCTTTGAAA 540
 518 AACAGCACGG GATCGATTTC TTTGAAGGG AATAAAATCGA GCGCAACAGG GAAAAAAGGT 600
 519 GGGGCTATT GTCTACTGG TACTGTAGAT ATTACAAATA ATACGGCTC TACCCCTTTC 660
 520 TCGAACATAA TTGCTGAAGC TCGAGGTGGA GCTATAAATA GCACAGGAACT CTGACAAATT 720
 521 ACAGGAAATA CGTCTCTGT ATTCTCTGAA ATATGTTGA CAGCGACCGC AGGAAATGGA 780
 522 GGAGCTCTTT CTGGAGATGC CGATGTTACG ATATCTGGGA ATCAGAGTGT AACTTTCTCA 840
 523 GGAAACCAAG CTGTAGCTA TGCGGAGGC ATTATGCTA AGAAGCTTAC ACTGGCTTCC 900
 524 GGGGGGGGGGG GGGGTATCTC CTTTCTAAC AATATGTTG AAGGTACAC TGCAGGTAAT 960
 525 GGTGAGGCC TTTCTATACT GGCGAGCTGG AAGTGTAGTC TTTCAGCAGA AGCAGGGGAC 1020
 526 ATTACCTTCA ATGGGAATGC CATTGTTGCA ACTACACAC AAACATACAAA AGAAATCT 1080
 527 ATTGACATAG GATCTACTGC AAAGATCAGC AATTACAGTG CAATATCTGG GCATAGCATC 1140
 528 TTTTCTACG ATCCGATTAC TGCTAACAGC ATTATGTTG CTTCTGAGTAC TTTAAATCTC 1200
 529 AATAAGGCTG ATGCAGGTAA TAGTACAGAT TATAGTGGG CGATTGTTT TTCTGGTGA 1260
 530 AAGCTCTCG AAGATGAAAGC AAAAGTTGCA GACAACCTCA CTTCTACGCT GAAGCAGCCT 1320
 531 GTAACTCTAA CTGAGGGAA TTAGTACTT AAACGTGGTG TCACTCTCGA TACGAAAGGC 1380
 532 TTTACTCAGA CCGCGGGTTC CTCTGTTATT ATGGATGCGG GCACAACGTT AAAAGCAAGT 1440
 533 ACAGAGGAGG TCACTTTAAC AGGTCTTCC ATTCTCTGAG ACTCTTTAGG CGAGGGTAAG 1500
 534 AAAGTGTAA TTGCTGCTTC TCCAGCAAGT AAAATGTAAG CCCTTAGTGG TCCGATCTT 1560
 535 CTTTTGGATA ACCAAGGGAA TCTTATGAA AATCACGACT TAGGAAAAAC TCAAGACTTT 1620
 536 TCATTGTC AGCTCTCTGC TCTGGTACT GCAACAACTA CAGATGTTCC AGGGGTCTC 1680
 537 ACAGTAGCAA CTCTTACGCA CTATGGTAT CAAGGACTT GGGGAATGAC TTGGGTGAT 1740
 538 GATACCGCAA GCACCTAAA GACTAAGACA GCGACATTAG CTTGGACCAA TACAGGCTAC 1800
 539 CTTCCGAATC CTGAGGCTCA AGGACCTTA GTTCTTAATA GCCTTGGGG ATCTTTTCA 1860
 540 GACATCCAAG CGATTCAAG TGTCATAGAG AGAAGTGTCT TGACTCTTTC TTCAAGATCGA 1920
 541 GGCTTCTGGG CTGCGGGACT CGCCAATTTC TTAGATAAAG ATAAGAAGG GGAAAACGC 1980
 542 AAATACCGTC ATAAATCTGG TGGATATGTC ATCGGAGGTG CAGCGCAAC TTGTTCTGAA 2040
 543 AACTTAATTG CTTTGCTTC TTGCAACTC TTTGGTAGCG ATAAGAATTG CTTAGTCCT 2100
 544 AAAATACATA CTGATACCTA TGAGGAGGC TTCTATATCC AACACATAC AGAATGTTG 2160
 545 GGGTCTAGT GTTGTCTCTT AGATAAACTT CCTGGCTCTT GGACTCATAA ACCCCCTCGT 2220
 546 TTAGAAGGGC AGCTCGCTTA TAGCCACGTC ACTAATGATC TGAAGACAAA GTATACTGCG 2280
 547 TATCTGAGG TGAAGGGTTC TTGGGGAAAT ATGCTTTA ACATGATGTT GGGAGCTTCT 2340
 548 TCTCATTCTT ATCCCTGAATA CCTGCATTTGTTGATAACCT ATGCTCCATA CATCAAACG 2400
 549 AATCTGACCT ATATACGCTA GGACAGCTTC TCGGAGAAAG GTACAGAAGG AAGATCTT 2460
 550 GATGACAGCA ACCTCTTCAA TTATCTTG CCTATAGGGG TGAAGTTGA GAACTCTCT 2520
 551 GATTGTAATG ACCTTTCTTA TGATCTGACT TTATCCTATG TTCCCTGATCT TATCCGAAT 2580
 552 GATCCAAAT GCACTACAGC ACTCTGAAATC AGCGGAGCCT TTGGGAAAC TTATGCCAAT 2640
 553 AACTTAGCAC GACAGGGCTT GCAAGTGGCT GCAGGGAGTC ACTACGCCCT CTCTCCCTATG 2700
 554 TTGAAAGTGC TCGGCCACTT TGTCTTGAA GTTCGTGGAT CCTCACGGAT TTATAATGTA 2760
 555 GATCTGGGG GTAAAGTCCA ATTCTAGGAG CGTCTCTCAT GTCTCAGAAA TTCTG 2815

743 (2) INFORMATION FOR SEQ ID NO: 5:

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/446,677

DATE: 09/06/2000
TIME: 11:01:37

Input Set : A:\Birkell.txt
Output Set: N:\CRF3\09062000\I44677.raw

745 (i) SEQUENCE CHARACTERISTICS:
746 (A) LENGTH: 3052 base pairs → SEQ P. 4
747 (B) TYPE: nucleic acid
748 (C) STRANDEDNESS: single
749 (D) TOPOLOGY: linear
751 (ii) MOLECULE TYPE: Genomic DNA
753 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
755 ATGCCATTTC CGCTCTGCGG ATTTCTCTA GTTTTTCTT TAACATTGCT CTCAGTCTTC 60
756 GACACTCTT TGAGTGTAC TACGATTTCT TAAACCCAG AAGATAGTT TCATGGAGAT 120
757 AGTCAGAATG CAGAACGTT TTATAATGTT CAAGCTGGG ATGCTCTAG CCTTACTGGT 180
758 GATGCTCAA TACTAACGT CGATAACTCT GCATTAATAA AAGCCTGCTT CAATGTGACC 240
759 TCAGGAAGTG TGAGCTTGC AGGAAATCAT CATGGTTAT ATTTAACAA TATTCCTCA 300
760 GGAATACAA AGGAAGGGC TGACTTTGT TGCAAGATC CTCAGCAAC GGCACGTTT 360
761 TCTGGTTCT CACCGCTTC TTTTATTCAAG AGCCCCGAG ATATTAAGA ACAGGGATGT 420
762 CTCTATTCAA AAAATGCACT TATGCTTTA ACAATTATG TAGTGCCTT TGAACAAAAC 480
763 CAAAGTAAGA CTAAGGCGG AGCTATTAGT GGGCGAATG TTACTATAGT AGGCAACTAC 540
764 GATTCCGTCT CTTCTATCA GAATGCAGCC ACTTTGGAG GTGCTATCCA TTCTTCAGGT 600
765 CCCCTACAGA TTGAGTAAAC TCAGGCAGAG ATAAGATTG CACAAATAC TGCCAAAGAAT 660
766 GTTCTGGAG GGGCTTGTG CTCCGATGTT GATATTGATA TTGATCAGAA TGCTTATGTT 720
767 CTATTCGAG AAAATGAGGC ATTGACTACT GCTATAGGTG AGGGAGGGC TGTCTGTG 780
768 CTTCCCACCT CAGGAAGTAG TACTCCAGTT CCTATTGTGA CTTTCTCTGA CAATAAACAG 840
769 TTAGTCTTG AAAGAACCA TTCCATAATG GGTGGCGAG CCATTATGTC TAGGAAACTT 900
770 AGCATCTCTT CAGGAGGTCC TACTCTATTT ATCAATAATA TATCATATGC AAATTGCAA 960
771 AATTAGGTG GAGCTATTGC CATTGATACT GGAGGGGAGA TCAGTTATC AGCAGAGAAA 1020
772 GGACAATTA CATTCCAAGG AAACCGGAGC AGCTTACCGT TTTTGAATGG CATCCATCTT 1080
773 TTACAAAATG CAAATTCTT GAAATTACAG GCGAGAAATG GATGCTCTAT AGAATTCTT 1140
774 GATCTTATTA CTCTGAAGC AGATGGGTCT ACCCAATGTA ATATCAACGG AGATCTAAA 1200
775 AATAAGAGT ACACAGGGC CATACTCTT TCTGGAGAAA AGAGTCTAGC AAACGATCCT 1260
776 AGGGATTTA ATCTACAACT CCCTCAGAAC GTCAACCTG CTCCAGGATA CTTAGTTATT 1320
777 AAAGAGGGG CCGAAGTCAC AGTTTCAAA TTCACGCAGT CTCCAGGATC GCATTAGTT 1380
778 TTAGATTTAG GAACCAAATC GATAGCCTCT AAGGAAGACA TTGCCATCAC AGGCCTCGCG 1440
779 ATAGATATAG ATAGCTTAAG CTCATCTCTA ACAGCAGCTG TTATTAAGC AAACACCGCA 1500
780 AATAAACAGA TATCCGTGAC GGACTCTATA GAACCTATCT CGCCTACTGG CAATGCCAT 1560
781 GAAGATCTCA GAATGAGAAA TTCACAGACG TTCCCTCTGC TCTCTTCTAGA GCCTGGAGCC 1620
782 GGGGTAGTG TGACTGTAAC TGCTGGAGAT TTCCCTACCGG TAAGTCCCCA TTATGGTTT 1680
783 CAAGGCAATT GGAAATTAGC TTGGACAGGA ACTGAAACA AAGTTGGAGA ATTCTTCTGG 1740
784 GATAAAATAA ATTATAAGCC TAGACCTGA AAAGAAGGAA ATTTAGTTCC TAATATCTTG 1800
785 TGGGGGAATG CTGTAATGT CAGATCCTTA ATGCAGGTTA AAGAGACCA TGCACTGAGC 1860
786 TTACAGACAG ATCGAGGGCT GTGGATCGAT GGAATTGGGA ATTTCTTCA TGTATCTGCC 1920
787 TCCGAAGACA ATATAAGGT ACGTCAAAAC AGCGGTGGAT ATGTTCTATC TGTAATAAT 1980
788 GAGATCACAC CTAAGCACTA TACTTCGATG GCATTTTCCC AACTCTTAG TAGAGACAAG 2040
789 GACTATGCGG TTTCACAAAC CAATACAGA ATGTTATTAG GATCGTATCT CTATCAATAT 2100
790 ACAACCTCCC TAGGAAATAT TTTCGTTAT GCTTCGGCTA ACCCTAATGT AAACGTCGGG 2160
791 ATTCCTCAA GAAGGTTCTC TCAAAATCTT CTTATGATT TTCATTTTT GTGCTTAT 2220
792 GGTATGCCA CCAATGATAT GAAAACAGAC TACGCAATT TCCTCTATGGT GAAAACAGC 2280
793 TGGAGAAACA ATTGTTGGGC TATAGACTGC GGAGGGAGCA TGCCCTTATT GGTTATTGAG 2340
794 AACGGAAGAC TTTTCCAAGG TGCCATCCCA TTATGAAAC TACAATTAGT TTATGCTTAT 2400
795 CAGGGAGATT TCAAAGAGAC GACTGCAGAT GCGCTAGAT TTAGTAATGG GAGTTAACCA 2460
796 TCGATTTCTG TACCTCTAGG CATACTGCTT GAGAAGCTGG CACTTCTCA GGATGTACTC 2520

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/446,677

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Input Set : A:\Birkell.txt
Output Set: N:\CRF3\09062000\I446677.raw

797 TATGACTTTA GTTTCTCCTA TATTCTGTAT ATTTCCGTA AGGATCCCTC ATGTGAAGCT
798 GCTCTGGTGA TTAGCGGAGA CTCCCTGGCTT GTTCCGGCAG CACACGTATC AAGACATGCT
799 TTTGTAGGGA GTGGACGGG TCGGTATCAC TTAACGACT ATACTGAGCT CTTATGTGCA
800 GGAAGTAGG AATGCCGCCC CCATGCTAGG AATTATAATA TAAACTGTGCG AAGCAAATT
801 CGTTTTAGA AGGTTCCAT TGCTGTGTC GTTCCGGATC TTAACATAA ATCCCTGACT
802 ATGGATCATA GGCATGGGT TTCTGAACT TGTGTGGAGA ATAACGACAT TTTATATGCA
803 TAACGGATA CTCGTATCAC CTCAGCCCCCT AGAGACATTC TTTAGGGTT CTTTATTTGTT
804 CTAAACTCG TATTTATCG AGAATCCTT ACGTTCTTGG TTTGCTTGTC TCCGAGGAGT
805 TCTCTAACGA ATCATAGGA TTCCAGGGTT CTGTTCCCTG AGTCCTTTG A

2580
2640
2700
2760
2820
2880
2940
3000
3052

E--> 2597 (2) INFORMATION FOR SEQ ID NO: 24:

2599 (i) SEQUENCE CHARACTERISTICS:

2600 (A) LENGTH: 946 amino acids → See p. 6
2601 (B) TYPE: amino acid
2602 (C) STRANDEDNESS: single
2603 (D) TOPOLOGY: linear

2605 (ii) MOLECULE TYPE: peptide

2607 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:

2609 Met Lys Thr Ser Val Ser Met Leu Leu Ala Leu Leu Cys Ser Gly Ala
2610 1 5 10 15
2611 Ser Ser Ile Val Leu His Ala Ala Thr Thr Pro Leu Asn Pro Glu Asp
2612 20 25 30
2613 Gly Phe Ile Gly Glu Gly Asn Thr Asn Thr Phe Ser Pro Lys Ser Thr
2614 35 40 45
2615 50 55 60
2616 65 70 75 80
2617 85 90 95
2618 100 105 110
2619 115 120 125
2620 130 135 140
2621 145 150 155 160
2622 165 170 175
2623 180 185 190
2624 195 200 205
2625 210 215 220
2626 225 230 235 240
2627 245 250 255
2628 265

Number of bases
conflict, 3052
listed, 3051 for

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/446,677

DATE: 09/06/2000
TIME: 11:01:38

Input Set : A:\Birkell.txt
Output Set: N:\CRF3\09062000\I446677.raw

2657 Ile Asn Cys Ser Gly Asp Leu Thr Phe Thr Asp Asn Thr Ser Leu Leu
2658 260 265 270
2660 Leu Gln Glu Asn Ser Thr Met Gln Asp Gly Gly Ala Leu Cys Ser Thr
2661 275 280 285
2663 Gly Thr Ile Ser Ile Thr Gly Ser Asp Ser Ile Asn Val Ile Gly Asn
2664 290 295 300
2666 Thr Ser Gly Gln Lys Gly Gly Ala Ile Ser Ala Ala Ser Leu Lys Ile
2667 305 310 315 320
2669 Leu Gly Gly Gln Gly Gly Ala Leu Phe Ser Asn Asn Val Val Thr His
2670 325 330 335
2672 Ala Thr Pro Leu Gly Gly Ala Ile Phe Ile Asn Thr Gly Ser Leu
2673 340 345 350
2675 Gln Leu Phe Thr Gln Gly Gly Asp Ile Val Phe Glu Gly Asn Gln Val
2676 355 360 365
2678 Thr Thr Ala Pro Asn Ala Thr Thr Lys Arg Asn Val Ile His Leu
2679 370 375 380
2681 Glu Ser Thr Ala Lys Trp Thr Gly Leu Ala Ala Ser Gln Gly Asn Ala
2682 385 390 395 400
2684 Ile Tyr Phe Tyr Asp Pro Ile Thr Thr Asn Asp Thr Gly Ala Ser Asp
2685 405 410 415
2687 Asn Leu Arg Ile Asn Glu Val Ser Ala Asn Gln Lys Leu Ser Gly Ser
2688 420 425 430
2690 Ile Val Phe Ser Gly Glu Arg Leu Ser Thr Ala Glu Ala Ile Ala Glu
2691 435 440 445
2693 Asn Leu Thr Ser Arg Ile Asn Gln Pro Val Thr Leu Val Glu Gly Ser
2694 450 455 460
2696 Leu Glu Leu Lys Gln Gly Val Thr Leu Ile Thr Gln Gly Phe Ser Gln
2697 465 470 475 480
2699 Glu Pro Glu Ser Thr Leu Leu Leu Asp Leu Gly Thr Ser Leu Gln Ala
2700 485 490 495
2702 Ser Thr Glu Asp Ile Val Ile Thr Asn Ser Ser Ile Asn Ala Asp Thr
2703 500 505 510
2705 Ile Tyr Gly Lys Asn Pro Ile Asn Ile Val Ala Ser Ala Ala Asn Lys
2706 515 520 525
2708 Asn Ile Thr Leu Thr Gly Thr Leu Ala Leu Val Asn Ala Asp Gly Ala
2709 530 535 540
2711 Leu Tyr Glu Asn His Thr Leu Gln Asp Ser Gln Asp Tyr Ser Phe Val
2712 545 550 555 560
2714 Lys Leu Ser Pro Gly Ala Gly Gly Thr Ile Ile Thr Gln Asp Ala Ser
2715 565 570 575
2717 Gln Lys Leu Leu Glu Val Ala Pro Ser Arg Pro His Tyr Gly Tyr Gln
2718 580 585 590
2720 Gly His Trp Asn Val Gln Val Ile Pro Gly Thr Gly Thr Gln Pro Ser
2721 595 600 605
2723 Gln Ala Asn Leu Glu Trp Val Arg Thr Gly Tyr Leu Pro Asn Pro Glu
2724 610 615 620
2726 Arg Gln Gly Phe Leu Val Pro Asn Ser Leu Trp Gly Ser Phe Val Asp
2727 625 630 635 640
2729 Gln Arg Ala Ile Gln Glu Ile Met Val Asn Ser Ser Gln Ile Leu Cys

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PATENT APPLICATION: US/09/446,677

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Input Set : A:\Birkell.txt
Output Set: N:\CRF3\09062000\I446677.raw

2730	645	650	655
2732	Gln Glu Arg Gly Val Trp Gly Ala Gly Ile Ala Asn Phe Leu His Arg	660	665
2733	660	665	670
2735	Asp Lys Ile Asn Glu His Gly Tyr Arg His Ser Gly Val Gly Tyr Leu	675	680
2736	675	680	685
2738	Val Gly Val Gly Thr His Ala Phe Ser Asp Ala Thr Ile Asn Ala Ala	690	695
2739	690	695	700
2741	Phe Cys Gln Leu Phe Ser Arg Asp Lys Asp Tyr Val Val Ser Lys Asn	705	710
2742	705	715	720
2744	His Gly Thr Ser Tyr Ser Gly Val Val Phe Leu Glu Asp Thr Leu Glu	725	730
2745	725	730	735
2747	Phe Arg Ser Pro Gln Gly Phe Tyr Thr Asp Ser Ser Ser Glu Ala Cys	740	745
2748	740	745	750
2750	Cys Asn Gln Val Val Thr Ile Asp Met Gln Leu Ser-Tyr Ser His Arg	755	760
2751	755	760	765
2753	Asn Asn Asp Met Lys Thr Lys Tyr Thr Thr Tyr Pro Glu Ala Gln Gly	770	775
2754	770	775	780
2756	Ser Trp Ala Asn Asp Val Phe Gly Leu Glu Phe Gly Ala Thr Thr Tyr	785	790
2759	785	790	795
2760	Tyr Tyr Pro Asn Ser Thr Phe Leu Phe Asp Tyr Tyr Ser Pro Phe Leu	805	810
2762	805	810	815
2763	Arg Leu Gln Cys Thr Tyr Ala His Gln Glu Asp Phe Lys Glu Thr Gly	820	825
2765	820	825	830
2766	Gly Glu Val Arg His Phe Thr Ser Gly Asp Leu Phe Asn Leu Ala Val	835	840
2768	835	840	845
2769	Pro Ile Gly Val Lys Phe Glu Arg Phe Ser Asp Cys Lys Arg Gly Ser	850	855
2771	850	855	860
2772	Tyr Glu Leu Thr Leu Ala Tyr Val Pro Asp Val Ile Arg Lys Asp Pro	865	870
2774	865	870	875
2775	Lys Ser Thr Ala Thr Leu Ala Ser Gly Ala Thr Trp Ser Thr His Gly	885	890
2777	885	890	895
2778	Asn Asn Leu Ser Arg Gln Gly Leu Gln Leu Arg Leu Gly Asn His Cys	900	905
2780	900	905	910
2781	Leu Ile Asn Pro Gly Ile Glu Val Phe Ser His Gly Ala Ile Glu Leu	915	920
2783	915	920	925
2784	Arg Gly Ser Ser Arg Asn Tyr Asn Ile Asn Leu Gly Gly Lys Tyr Arg	930	935
2786	930	940	945

E--> 2787 945 → 945 amino acids found, 946 listed as length.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/446,677

DATE: 09/06/2000
TIME: 11:01:39

Input Set : A:\Birkelli.txt
Output Set: N:\CRF3\09062000\1446677.raw

L:31 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:32 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:38 M:220 C: Keyword misspelled or invalid format, [(vii) PRIOR APPLICATION DATA:]
L:510 M:320 E: (1) Wrong Nucleic Acid Designator, 2
L:805 M:254 E: No. of Bases conflict, Input:3052 Counted:3051 SEQ:5
L:805 M:204 E: No. of Bases differ, LENGTH:Input:3052 Counted:3051 SEQ:5
L:2454 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:2787 M:203 E: No. of Seq. differs, LENGTH:Input:946 Found:945 SEQ:24